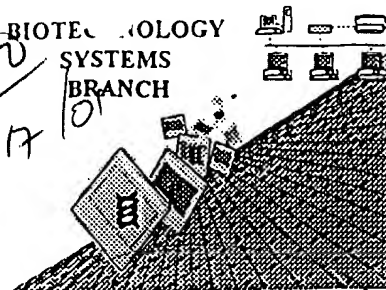


## RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY  
SYSTEMS  
BRANCH

7500  
10.17.01

0420/03cc



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/682,769

Source: OIPE

Date Processed by STIC: 10/18/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER  
VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND  
TRADEMARK OFFICE WEBSITE. SEE BELOW:

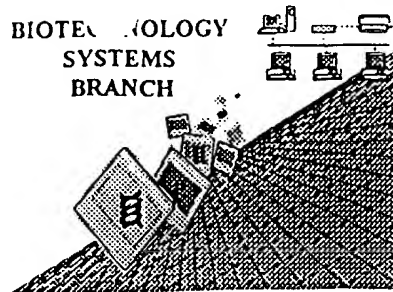
### Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>



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<http://www.uspto.gov/web/offices/pac/checker>

CRF  
OIPE

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/682,769

DATE: 10/18/2001

TIME: 06:50:00

Input Set : C:\Crf3\Datahold\EFS\09682769\PTO\_MS.txt

Output Set: N:\CRF3\10182001\I682769.raw

1 <110> APPLICANT: Monsanto Co Rangwala,  
W--> 2 <120> TITLE OF INVENTION: Cotton Event PV-GHGT07(1445) and Compositions and  
W--> 3 <130> FILE REFERENCE: 38-21(52249)<150> 60/243190<151> 2000-10-  
C--> 4 <140> CURRENT APPLICATION NUMBER: US/09/682,769  
C--> 4 <141> CURRENT FILING DATE: 2001-10-17  
E--> 4 <160> NUMBER OF SEQ ID: 12 <170> PatentIn version 3.0<210> 1<211> 25<212> DNA<213>

## ERRORED SEQUENCES

Does Not Comply  
Corrected Diskette Needed

See print out of the Sequence Listing  
attached to this report. PatentIn failed  
to save the file in the correct format.  
Please contact Bob Wax at 703-308-4214  
for help in getting file in correct format.

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/682,769

DATE: 10/18/2001

TIME: 06:50:01

Input Set : C:\Crf3\Datahold\EFS\09682769\PTO\_MS.txt

Output Set: N:\CRF3\10182001\I682769.raw

L:2 M:283 W: Missing Blank Line separator, <120> field identifier  
L:3 M:283 W: Missing Blank Line separator, <130> field identifier  
L:4 M:270 C: Current Application Number differs, Replaced Current Application No  
L:4 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:4 M:283 W: Missing Blank Line separator, <160> field identifier  
L:5 M:282 W: Numeric Field Identifier Missing, <210> is required.  
L:5 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:5 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:5 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:5 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO  
L:5 M:283 W: Missing Blank Line separator, <400> field identifier  
L:24 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:24 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:24 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:31 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:31 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:31 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:38 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:38 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:38 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:55 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:55 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:55 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:78 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:78 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:78 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:85 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:85 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:85 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:92 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:92 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:92 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:99 M:282 W: Numeric Field Identifier Missing, <211> is required.  
L:99 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:99 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:4 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (12) Counted (9)